OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/642,405

DATE: 07/05/2001 TIME: 16:38:23

Input Set : A:\20413y.txt

```
4 <110> APPLICANT: Neeper, Michael P.
                                                               ENTERED
              McClements, William L.
              Jansen, Kathrin U.
              Schultz, Loren D.
              Chen, Ling
              Wang, Xin-Min
     11 <120> TITLE OF INVENTION: SYNTHETIC HUMAN PAPILLOMAVIRUS GENES
     14 <130> FILE REFERENCE: 20413Y
     16 <140> CURRENT APPLICATION NUMBER: 09/642,405
C--> 17 <141> CURRENT FILING DATE: 2001-06-21
     19 <150> PRIOR APPLICATION NUMBER: PCT/US00/22932
     20 <151> PRIOR FILING DATE: 2000-08-21
     22 <150> PRIOR APPLICATION NUMBER: 60/210,143
    23 <151> PRIOR FILING DATE: 2000-06-07
    25 <150> PRIOR APPLICATION NUMBER: 60/150,728
    26 <151> PRIOR FILING DATE: 1999-08-25
    28 <160> NUMBER OF SEQ ID NOS: 150
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    33 <211> LENGTH: 1518
    34 <212> TYPE: DNA
    35 <213> ORGANISM: Artificial Sequence
    37 <220> FEATURE:
    38 <223> OTHER INFORMATION: Codon-Optimized HPV16 L1
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    42 gtggtgagca ccgacgagta cgtggcccgc accaacatct actaccacgc cggcaccagc
                                                                              120
    43 cgcctgctgg ccgtgggcca cccctacttc cccatcaaga agcccaacaa caacaagatc
                                                                              180
    44 ctggtgccca aggtgagcgg cctgcagtac cgcgtgttcc gcatccacct gcccgacccc
                                                                              240
    45 aacaagttcg gcttccccga cacaagcttc tacaaccccg acacccagcg cctggtgtgg
                                                                              300
    46 gcctgcgtgg gcgtggaggt gggccgcggc cagcccctgg gcgtgggcat cagcggccac
                                                                              360
    47 cccctgctga acaagctgga cgacaccgag aacgccagcg cctacgccgc caacgccggc
                                                                              420
    48 gtggacaacc gcgagtgcat cagcatggac tacaagcaga cccagctgtg cctgatcggc
                                                                              480
    49 tgcaagcete ccateggega gcactgggge aagggcagee cctgcaccaa cgtggcegtg
                                                                              540
    50 aacceeggeg actgeeetee eetggagetg atcaacaceg tgateeagga eggegacatg
                                                                              600
    51 gtggacaccg gcttcggcgc catggacttc accaccctgc aggccaacaa gagcgaggtg
                                                                              660
    52 cccctggaca tctgcaccag catctgcaag taccccgact acatcaagat ggtgagcgag
                                                                             720
    53 ccctacggcg acagcctgtt cttctacctg cgccgcgagc agatgttcgt gcgccacctg
                                                                             780
   54 ttcaaccgcg ccggcgccgt gggcgagaac gtgcccgacg acctgtacat caagggcagc
                                                                             840
   55 ggcagcaccg ccaacctggc cagcagcaac tacttcccca ctcccagcgg cagcatggtg
                                                                             900
   56 accagegacg cecaaatett caacaagece tactggetge agegegecea gggecacaae
                                                                             960
   57 aacggcatct gctggggcaa ccagctgttc gtgaccgtgg tggacaccac ccgcagcacc
                                                                            1020
   58 aacatgagee tgtgegeege cateageace agegagacea cetacaagaa caccaactte
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   59 aaggagtace tgegeeacgg egaggagtac gacetgeagt teatetteea getgtgeaag
                                                                            1140
   60 atcaccetga cegeegaegt gatgacetae atceaeagea tgaacageae catcetggag
                                                                            1200
   61 gactggaact teggeetgea geeeecteee ggeggtaeee tggaggaeae etaeegette
                                                                            1260
   62 gtgaccagee aggeeatege etgeeagaag cacaccete eegeteecaa ggaggateee
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Input Set : A:\20413y.txt

| 63 ctgaagaagt acacettetg ggaggtgaac etgaaggaga agtteagege egacetggae | 1380 |
|---|------|
| 64 cagttccccc tgggccgcaa gttcctgctg caggccggcc tgaaggccaa gcccaagttc | 1440 |
| 65 accetgggea agegeaagge caceeceace accageagea ecageaecae egeeaagege | 1500 |
| 66 aagaagcgca agctgtaa | 1518 |
| 68 <210> SEQ ID NO: 2 | |
| 69 <211> LENGTH: 1950 | |
| 70 <212> TYPE: DNA | |
| 71 <213> ORGANISM: Artificial Sequence | |
| 73 <220> FEATURE: | |
| 74 <223> OTHER INFORMATION: Mutant, Codon-Optimized HPV16 E1 | |
| 76 <400> SEQUENCE: 2 | |
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| 78 gtggaggccg tggtggagaa gaagaccggc gacgccatca gcgacgacga gaacgagaac | 120 |
| 79 gacagegaca eeggegagga eetggtggae tteategtga aegacaaega etaeetgaee | 180 |
| 80 caggeegaga eegagaeege eeaegeeetg tteaeegeee aggaggeeaa geageaeege | 240 |
| 81 gacgccgtgc aggtgctgaa gcgcaagtac ctgggcagcc ccctgagcga catcagcggc | 300 |
| 82 tgcgtcgaca acaacatcag cccccgcctg aaggccatct gcatcgagaa gcagagccgc | 360 |
| 83 gccgccaagc gccgcctgtt cgagagcgag gacagcggct acggcaacac cgaggtggag | 420 |
| 84 acccagcaga tgctgcaggt ggagggccgc cacgagaccg agaccccctg cagccagtac | 480 |
| 85 ageggeggea geggeggegg etgeageeag tacageageg geageggegg egagggegtg | 540 |
| 86 agcgagcgcc acaccatctg ccagacccct ctgaccaaca tcctgaacgt gctgaagacc | 600 |
| 87 agcaacgcca aggccgccat gctggccaag ttcaaggagc tgtacggcgt gagcttcagc | 660 |
| 88 gagetggtge geceetteaa gageaacaag ageaeetget gegaetggtg categeegee | 720 |
| 89 ttcggcctga cccccagcat cgccgacagc atcaagaccc tgctgcagca gtactgcctg | 780 |
| 90 tacctgcaca tccagagect ggeetgeage tggggeatgg tggtgetget getggtgege | 840 |
| 91 tacaagtgcg gcaagaaccg cgagaccatc gagaagctgc tgagcaagct gctgtgcgtg | 900 |
| 92 agececatgt geatgatgat egageeteee aagettegea geacegeege egecetgtae | 960 |
| 93 tggtacaaga ccggcatcag caacatcagc gaggtgtacg gcgacacccc cgagtggatc | 1020 |
| 94 cagcgccaga ccgtgctgca gcacagcttc aacgactgca ccttcgagct gagccagatg | 1080 |
| 95 gtgcagtggg cctacgacaa cgacatcgtg gacgacagcg agatcgccta caagtacgcc | 1140 |
| 96 cagetggeeg acaccaacag caacgecage geetteetga agageaacag ceaggeeaag | 1200 |
| 97 atcgtgaagg actgcgccac catgtgccgc cactacaagc gcgccgagaa gaagcagatg | 1260 |
| 98 agcatgagee agtggateaa gtacegetge gaeegegtgg acgaeggegg egaeegeaag | 1320 |
| 99 cagatogtga tgttootgog ctaccagggo gtggaattoa tgagottoot gaccgoootg | 1380 |
| 100 aagcgcttcc tgcagggcat ccccaagaag aactgcatcc tgctgtacgg cgccgccaac | 1440 |
| 101 accgacaaga geetgttegg catgageetg atgaagttee tgeagggeag egtgatetge | 1500 |
| 102 ttcgtgaaca gcaagagcca cttctggctg cagcccctgg ccgacgccaa gatcggcatg | 1560 |
| 103 ctggacgacg ccaccgtgcc ctgctggaac tacatcgacg acaacctgcg caacgccctg | 1620 |
| 104 gacggcaacc tggtgagcat ggacgtgaag caccgccccc tggtgcagct gaagtgccct | 1680 |
| 105 cccctgctga tcaccagcaa catcaacgcc ggcaccgaca gccgctggcc ctacctgcac | 1740 |
| 106 aaccgcctgg tggtgttcac cttccccaac gagttcccct tcgacgagaa cggtaacccc | 1800 |
| 107 gtgtacgagc tgaacgacaa gaactggaag agcttcttca gccgcacctg gagccgcctg | 1860 |
| 108 agectgeacg aggacgagga caaggagaac gacggegaea geetgeeeac etteaagtge | 1920 |
| 109 gtgagcggcc agaacaccaa caccctgtaa | 1950 |
| 111 <210> SEQ ID NO: 3 | |
| 112 <211> LENGTH: 1098 | |
| 113 <212> TYPE: DNA | |
| 114 <213> ORGANISM: Artificial Sequence | |
| 116 <220> FEATURE: | |

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DATE: 07/05/2001 PATENT APPLICATION: US/09/642,405 TIME: 16:38:23

Input Set : A:\20413y.txt

| 117 .000 | |
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| 117 <223> OTHER INFORMATION: Mutant, Codon-Optimized HPV16 E2 | |
| 119 <400> SEQUENCE: 3 | |
| 120 atggagacce tgtgccagcg cetgaacgtg tgccaggaca agatectgac ccactacgag | 60 |
| The dadydcayca coyacciqcy cyaccacate gaetactaga aggagataga agtarati | 120 |
| 122 goodcoldot doddydcoog cydgalagac ffcaaggaga fgaaggaga gafaaf | 180 |
| 120 decetygety tydytadydd Cddygeerf caggerraeg agetgaagat gaagat ar | 240 |
| 121 decaretata acayecayta cagcaacgaa aagtggacge tagaagagat gagaat | 300 |
| 123 gegeneerya cogoocodo cogotigoato aagaaggagg gotagagget googet a | 360 |
| 120 coogacygcy acatolycad caccatgoac tacaccaact gracogacat gtacatata | 420 |
| 127 gaggaggeea gegradeedt agtggaggge caggtggaet actaoggeet gtaotaget | 480 |
| 120 cacyayyyca teegeaceta etteatacaa tteaaagaaca acacaaaaa ataaaaa | 540 |
| 123 dacadygryr gygdygrgca cgccggcggc caggtgatcc tgtgggggg gagart | 600 |
| Too ageageage aggregated coccagadace afroncease acetagagaa aggregate | 660 |
| 131 geodeedda eeddggeegt ggeetggge accgagagaa cegagagaa cataaa | 720 |
| 132 deceyeageg ageoegaede eggeaaceee torrarara coaagatast garages | 780 |
| 133 agegeggaea gegeeecat ectgaecage tteaacagea gagaeaaga | 840 |
| 191 egeddedged dedeedeec eategtgeac etgaaggggg acggaaaga gataraat | 900 |
| 100 orgegorace gericadydd geaergeang etgraegaega gogtangan an 110 | 960 |
| to cygaccyyco acadequaa qeacaagage geeategtga coetgagota con ca | 1020 |
| To eggedgegeg accayticet dadecaddia aagateeeca agaceatooo agtgoggaa | 1020 |
| 130 ggcccatga qcatctaa | 1098 |
| 140 <210> SEQ ID NO: 4 | 1090 |
| 141 <211> LENGTH: 297 | |
| 142 <212> TYPE: DNA | |
| 143 <213> ORGANISM: Artificial Sequence | |
| 145 <220> FEATURE: | |
| 146 <223> OTHER INFORMATION: Mutant, Codon-Optimized HPV16 E7 | |
| 140 /400/ SEQUENCE: 4 | |
| 149 atgeacggeg acacececae cetgeacgag tacatgetgg acetgeagee egagaceaec | 60 |
| Too gaccegeacy gotacggcca gctgaacgac agcagcgaag aggaggagaga | |
| 101 Cocyclyged addecdaded edaceacacacacacacacacacacacacacacacacaca | 120 |
| dydyddagda ddelgdgddi gladdigdag agcadddag fgaacataga cacaataga | 180 |
| and a successful and a | 240 |
| 193 (510) SEG ID NO: 2 | 297 |
| 156 <211> LENGTH: 297 | |
| 157 <212> TYPE: DNA | |
| 158 <213> ORGANISM: Artificial Sequence | |
| 160 <220> FEATURE: | |
| 161 <223> OTHER INFORMATION: Codon-Optimized HPV6a E7 | |
| 163 <400> SEQUENCE: 5 | |
| 164 atgcacggec gecacgtgac cetgaaggac ategtgetgg acetgeagec teeegaceee | 60 |
| a supplied a contraction of the | 60 |
| gacygecayy acayecayee cetgaagcag cacttegaga togtgaagtg starta | 120 |
| To design action action and the state of the | 180 |
| saysteetige tygglactet gdatateara faceceatet acaatagaaa anaata- | 240 |
| 1,0 /510/ 2EG ID MO: 0 | 297 |
| 171 <211> LENGTH: 318 | |
| 172 <212> TYPE: DNA | |
| 173 <213> ORGANISM: Artificial Sequence | |
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RAW SEQUENCE LISTING DATE: 07/05/2001 PATENT APPLICATION: US/09/642,405 TIME: 16:38:23

Input Set : A:\20413y.txt

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| 175 | <220> FEATURE: | | | | | |
|-----|-----------------------|--------------|----------------|------------|------------|------|
| 176 | <223> OTHER INFORMATI | ON: Codon-O | ptimized HPV | √18 E7 | | |
| 178 | <400> SEQUENCE: 6 | | | | | |
| 179 | atgcacggcc ccaaggccac | cctgcaggac | atcgtgctgc | acctggagcc | ccagaacgag | 60 |
| 180 | atccccgtgg acctgctgtg | ccacgagcag | ctgagcgaca | gcgaggagga | gaacgacgag | 120 |
| 181 | atcgacggcg tgaaccacca | gcacctgccc | gctcgcaggg | ccgagcccca | gcgccacacc | 180 |
| 182 | atgctgtgca tgtgctgcaa | gtgcgaggcc | cgcatcgagc | tggtggtgga | gagcagcgct | 240 |
| | gacgacetge gegeetteca | | | | | 300 |
| | tgcgccagcc agcagtaa | | - | | | 318 |
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| | <211> LENGTH: 1107 | | | | | |
| | <212> TYPE: DNA | | | | | |
| 189 | <213> ORGANISM: Artif | icial Seque | nce | | | |
| | <220> FEATURE: | _ | | | | |
| | <223> OTHER INFORMATI | ON: Codon-O | otimized HPV | 76a E2 | | |
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| | gagaacagca ccgacctgca | | | | | 120 |
| | gtgctgctgt acaaggccaa | | | - | | 180 |
| | cctctgaagg tgagcgaggc | | | | | 240 |
| | agcctgctgc gcaccgagta | | | | | 300 |
| | atgtggcaga ccctcccaa | | | | | 360 |
| | ttcgacqqct qcqccaacaa | | | | | 420 |
| | gacaacgaca cctgggtgaa | | | | | 480 |
| | tqtqqccaqt tcaaqaccta | | | | | 540 |
| | accaagcact gggaggtgtg | | | | | 600 |
| | agcaccaccc aggaggtgag | | | | | 660 |
| | accetqqtqa qcaqcagcac | | | | | 720 |
| | cgcggcgtgc agcagagccc | | | | | 780 |
| | agcggcaacc acaacctgat | | | | | 840 |
| | aacagcagcg ccactcccat | | | | | 900 |
| | cgctaccgcc tgaacgatcg | | | | | 960 |
| | tgggccagca gcaaggctcc | | | | | 1020 |
| | gagcagcqcc agcagttcct | | | | | 1080 |
| | ggcttcatga gcctgcacct | | aagacccccc | ccaccaccag | coacaageeg | 1107 |
| | <210> SEQ ID NO: 8 | geegeaa | | | | 1107 |
| | <211> LENGTH: 1098 | | | | | |
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| | <213> ORGANISM: Artif | icial Seque | nce | | | |
| | <220> FEATURE: | iciai ocquei | | | | |
| | <223> OTHER INFORMATI | ON: Codon-Or | ntimized HDV | 718 F2 | | |
| | <400> SEQUENCE: 8 | on. codon o | Jeimizea III (| TO EZ | | |
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| | qaccactacq aqaacqacaq | | | | | 120 |
| | cgctgggaga acgccatctt | | | | | 180 |
| | caggtggtgc ccgcctacaa | | | | | 240 |
| | atggccctgc agggcctggc | | | | | 300 |
| | acctgcgagg agctgtggaa | | | | | 360 |
| | gtgcaggtgt acttcgacgg | | | | | 420 |
| 230 | grycaggryr accregacyg | caacaayyac | auctycatya | accacycyge | cegggacage | 420 |

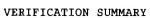
RAW SEQUENCE LISTING

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Input Set : A:\20413y.txt

| 231 | gtgtactaca tgaccgacgc cggcacctgg gacaagaccg ccacctgcgt gagccaccgc | 480 |
|-----|--|------------|
| 232 | ggcctgtact acgtgaagga gggctacaac accttctaca tcgagttcaa gagcgagtgc | 540 |
| 233 | gagaagtacg gcaacaccgg cacctgggag gtgcacttcg gcaacaacgt gatcgactge | 600 |
| 234 | aacgacagca tgtgcagcac cagcgacgac accgtgagcg ccacccagct ggtgaagcag | 660 |
| 235 | ctgcagcaca ctcccagccc ctacagcagc accgtgagcg tgggcaccgc caagacctac | 720 |
| 236 | ggccagacca gcgccgccac tcgccctqqc cactqcgqcc tggccgagaa gcagcactgc | 780 |
| 237 | gggccgtga accetetget gggcgccgcc accqccaccg gcaacaacaa gcgccgcaag | 840 |
| 238 | ctatacagea geaacaceae teccateate caeetgaaga gegaeegeaa cageetgaag | 900 |
| 239 | tracetricing according canaderacade daccactace degacatead cadeacetrag | 960 |
| 240 | cactggaccg gcgccgggaa cgagaagacc ggcatcctga ccgtgaccta ccacagcgag | 1020 |
| 241 | acccagegca ccaagttcct gaacaccgtg gccatececg acagegtgca gatectggtg | 1080 |
| 242 | ggctacatga ccatgtaa | 1098 |
| | <210> SEQ ID NO: 9 | |
| | <211> LENGTH: 129 | |
| | <212> TYPE: DNA | |
| 247 | <213> ORGANISM: Artificial Sequence | |
| 249 | <220> FEATURE: | |
| 250 | <223> OTHER INFORMATION: Codon-Optimized HPV16 L1 fragment | |
| 252 | <400> SEQUENCE: 9 | C 0 |
| 253 | atgageetgt ggetgeecag egaggeeace gtgtaeetge etecegtgee egtgageaag | 60 |
| 254 | gtggtgagca ccgacgagta cgtggcccgc accaacatet actaccacge cggcaccage | 120 |
| | cgcctgctg | 129 |
| 257 | <210> SEQ ID NO: 10 | |
| 258 | <211> LENGTH: 129 | |
| | <212> TYPE: DNA | |
| | <213> ORGANISM: Artificial Sequence | |
| 262 | <220> FEATURE: | |
| | <223> OTHER INFORMATION: Codon-Optimized HPV16 L1 fragment | |
| 265 | <400> SEQUENCE: 10 | 60 |
| 266 | cgcatccacc tgcccgaccc caacaagttc ggcttccccg acacaagctt ctacaacccc | 120 |
| 267 | gacacccage geetggtgt ggeetgegtg ggegtggagg tgggeegegg ceageccetg | 120 |
| | ggcgtgggc | 129 |
| | <210> SEQ ID NO: 11 | |
| | <211> LENGTH: 129 | |
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| | <213> ORGANISM: Artificial Sequence | |
| 275 | <220> FEATURE: | |
| | <223> OTHER INFORMATION: Codon-Optimized HPV16 L1 fragment | |
| 278 | <400> SEQUENCE: 11 | 60 |
| 279 | gagtgeatea geatggaeta caageagaee cagetgtgee tgateggetg caageeteee | 120 |
| | ateggegage actggggeaa gggeageeee tgcaccaacg tggeegtgaa eeeeggegae | 129 |
| | tgccctccc | |
| | <210> SEQ ID NO: 12 | |
| | <211> LENGTH: 132 | |
| | <212> TYPE: DNA | |
| | <213> ORGANISM: Artificial Sequence | |
| 288 | <pre><220> FEATURE: <223> OTHER INFORMATION: Codon-Optimized HPV16 L1 fragment</pre> | |
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| 291 | <400> SEQUENCE: 12 | |
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PATENT APPLICATION: US/09/642,405

DATE: 07/05/2001

TIME: 16:38:24

Input Set : A:\20413y.txt
Output Set: N:\CRF3\07032001\I642405.raw

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date